#### **TRANSMITTAL**

Applicant

Stordeur, et al.

App. No.

10/501,666

Filed

July 16, 2004

For

METHOD TO DETERMINE

IN VIVO NUCLEIC ACID

**LEVELS** 

Examiner

Unknown

Art Unit

Unknown

#### **CERTIFICATE OF MAILING**

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December 20, 2004

(Date)

Che Swyden Chereskin, Ph.D., Reg. No. 41,466

Commissioner for Patents P.O. Box 1450 Alexandria, VA 22313-1450

Sir:

Transmitted herewith for filing in the above-identified application are the following enclosures:

- Response to Notice to Comply in 3 pages. (X)
- (X) Paper Sequence Listing in 23 pages and two (2) CRF diskettes containing same.
- Return prepaid postcard. (X)
- (X) Please charge any additional fees, including any fees for additional extension of time, or credit overpayment to Deposit Account No. 11-1410.

Che Swyden Chereskin, Ph.D.

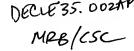
Registration No. 41,466

Agent of Record

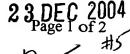
Customer No. 20,995

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U.S. APPLICATION NUMBER NO.

FIRST NAMED APPLICANT

ATTY. DOCKET NO.

10/501,666

Patrick Stordeur

DECLE35.002APC

INTERNATIONAL APPLICATION NO.

PCT/EP03/00493

I.A. FILING DATE

PRIORITY DATE

01/20/2003

01/18/2002

20995 KNOBBE MARTENS OLSON & BEAR LLP 2040 MAIN STREET FOURTEENTH FLOOR **IRVINE, CA 92614** 



**CONFIRMATION NO. 9946 371 FORMALITIES LETTER** \*OC00000014657750\*

Date Mailed: 12/03/2004

## NOTIFICATION TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE AND/OR AMINO ACID SEQUENCE DISCLOSURES

Applicant is given TWO MONTHS FROM THE DATE OF THIS NOTICE within which to file the items indicated below to avoid abandonment. Extensions of time may be obtained under the provisions of 37 CFR 1.136(a).

The following items MUST be furnished within the period set forth below:

- The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 CFR 1.821-1.825 for the following reason(s):
  - Diskette does not comply. A corrected diskette is required. See the attached raw sequence listing error report.
  - APPLICANT MUST PROVIDE:
    - An initial or substitute paper copy or compact disc of the "Sequence Listing," as well as an amendment directing its entry into the specification.
- For questions regarding compliance to 37 CFR 1.821-1.825 requirements, please contact:
  - For Rules Interpretation, call (703) 308-4216
  - To Purchase Patentin Software, call (703) 306-2600
  - For Patentln Software Program Help, call (703) 306-4119 or e-mail at patin21help@uspto.gov or patin3help@uspto.gov

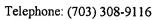
Additionally the following defects have been observed:

Annexes have not been entered because the applicant does not want the annexes entered..

Applicant is reminded that any communications to the United States Patent and Trademark Office must be mailed to the address given in the heading and include the U.S. application no. shown above (37 CFR 1.5)

A copy of this notice MUST be returned with the response.

JOHN L ANDERSON



## PART 1 - ATTORNEY/APPLICANT COPY

U.S. APPLICATION NUMBER NO.	INTERNATIONAL APPLICATION NO.	ATTY. DOCKET NO.
10/501,666	PCT/EP03/00493	DECLE35.002APC

FORM PCT/DO/EO/922 (371 Formalities Notice)

\* Send with Form

DEC 2 3 2004

## RAW SEQUENCE LISTING RADEMARKET ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	10/501,666
Source:	1.47/10
Date Processed by STIC:	7/23/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

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http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (<http://www.uspto.gov/ebc/efs/downloads/documents.htm>, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
- Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
   U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lol Room 1B03, Arlington, VA 22202

Revised 05/17/04



RAW SEQUENCE LISTING DATE: 07/23/2004 PATENT APPLICATION: US/10/501,666 TIME: 15:25:18

Input Set : A:\seq lst US.txt

```
5 <110> APPLICANT: Stordeur, Patrick
               Goldman, Michel
     10 <120> TITLE OF INVENTION: Method to determine in vivo nucleic acid levels
     14 <130> FILE REFERENCE: DECLE35.002APC
C--> 17 <140> CURRENT APPLICATION NUMBER: US/10/501,666
C--> 18 <141> CURRENT FILING DATE: 2004-07-16
                                                                   Does Not Comply
     21 <150> PRIOR APPLICATION NUMBER: PCT/EP03/00493
     22 <151> PRIOR FILING DATE: 2003-01-20
     24 <150> PRIOR APPLICATION NUMBER: EP 02447009.8
     26 <151> PRIOR FILING DATE: 2002-01-18
     30 <160> NUMBER OF SEQ ID NOS: 55
     34 <170> SOFTWARE: PatentIn version 3.1
                                                                    Corrected Dickette Needed
     38 <210> SEQ ID NO: 1
     40 <211> LENGTH: 21
     42 <212> TYPE: DNA
     44 <213> ORGANISM: Artificial Sequence
     48 <220> FEATURE:
     50 <223> OTHER INFORMATION: Oligonucleotide
     52 <400> SEQUENCE: 1
     53 gaagatgtgc ctgtcctgtg t
                                                                                    21
     56 <210> SEQ ID NO: 2
     58 <211> LENGTH: 21
     60 <212> TYPE: DNA
     62 <213> ORGANISM: Artificial Sequence
     66 <220> FEATURE:
     68 <223> OTHER INFORMATION: Oligonucleotide
     70 <400> SEQUENCE: 2
     71 cgctcaggtc agtgatgtta a
                                                                                   21
     74 <210> SEQ ID NO: 3
     76 <211> LENGTH: 27
     78 <212> TYPE: DNA
     80 <213> ORGANISM: Artificial Sequence
     84 <220> FEATURE:
 86 <223> OTHER INFORMATION: Oligonucleotide
    88 <220> FEATURE:
                                                n'can only represent a single revolection,
nothing else. Suggestion: in

the 22207-22237 sections,

indicate that the first and

last newlitides are labelled

last muliotides are labelled
     90 <221> NAME/KEY: misc_feature
     92 <222> LOCATION: (1)..(1)
    94 <223> OTHER INFORMATION N = 6Fam
    98 <220> FEATURE:
    100 <221> NAME/KEY: misc_feature
    102 <222> LOCATION: (27)..(27)
    104 <223 > OTHER INFORMATION: N = Tamra-p
    108 <400> SEQUENCE: 3
```

DATE: 07/23/2004 PATENT APPLICATION: US/10/501,666 TIME: 15:25:18

Input Set : A:\seq\_lst\_US.txt

```
W--> 109 ntggtgatga gaccagactc cagctgn
                                                                               27
       112 <210> SEQ ID NO: 4
       114 <211> LENGTH: 21
       116 <212> TYPE: DNA
       118 <213> ORGANISM: Artificial Sequence
       122 <220> FEATURE:
       124 <223> OTHER INFORMATION: Oligonucleotide
       126 <400> SEQUENCE: 4
       127 acagatgaag tgctccttcc a
                                                                               21
       130 <210> SEQ ID NO: 5
       132 <211> LENGTH: 21
       134 <212> TYPE: DNA
       136 <213 > ORGANISM: Artificial Sequence
       140 <220> FEATURE:
       142 <223> OTHER INFORMATION: Oligonucleotide
       144 <400> SEQUENCE: 5
       145 gtcggagatt cgtagctgga t
                                                                               21
       148 <210> SEQ ID NO: 6
       150 <211> LENGTH: 22
       152 <212> TYPE: DNA
      154 <213> ORGANISM: Artificial Sequence
      158 <220> FEATURE:
      160 <223> OTHER INFORMATION: Oligonucleotide
      162 <220> FEATURE:
      164 <221> NAME/KEY: misc_feature
      166 <222> LOCATION: (1)..(1)
      168 <223> OTHER INFORMATION: N = 6Fam
                                                same ever as page I
      172 <220> FEATURE:
      174 <221> NAME/KEY: misc_feature
      176 <222> LOCATION: (22)..(22)
      178 <223> OTHER INFORMATION: N = Tamra-p
      182 <400> SEQUENCE: 6
 W--> 183 netetgeeet etggatggeg gn
                                                                               22
      186 <210> SEQ ID NO: 7
      188 <211> LENGTH: 20
      190 <212> TYPE: DNA
      192 <213> ORGANISM: Artificial Sequence
      196 <220> FEATURE:
      198 <223> OTHER INFORMATION: Oligonucleotide
      200 <400> SEQUENCE: 7
      201 agctgcctac gtgtatgcca
                                                                               20
      204 <210> SEQ ID NO: 8
206 <211> LENGTH: 21
                             208 <212> TYPE: DNA
      210 <213> ORGANISM: Artificial Sequence
      214 <220> FEATURE:
      216 <223> OTHER INFORMATION: Oligonucleotide
      218 <400> SEQUENCE: 8
      219 gcagtgccaa ggtctctttc a
                                                                              21
```

PATENT APPLICATION: US/10/501,666

DATE: 07/23/2004 TIME: 15:25:18

Input Set : A:\seq\_lst US.txt

Output Set: N:\CRF4\07232004\J501666.raw

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222 <210> SEQ ID NO: 9
     224 <211> LENGTH: 29
     226 <212> TYPE: DNA
     228 <213> ORGANISM: Artificial Sequence
    232 <220> FEATURE:
     234 <223> OTHER INFORMATION: Oligonucleotide
    236 <220> FEATURE:
    238 <221> NAME/KEY: misc feature
    240 <222> LOCATION: (1)..(1)
    242 <223 > OTHER INFORMATION: N = 6Fam
    246 <220> FEATURE:
                                                 same
    248 <221> NAME/KEY: misc_feature
    250 <222> LOCATION: (29)..(29)
    252 <223> OTHER INFORMATION: N = Tamra-p
    256 <400> SEQUENCE: 9
W--> 257 nccccacaga aattcccaca agtgcattn
                                                                                 29
    260 <210> SEQ ID NO: 10
    262 <211> LENGTH: 21
    264 <212> TYPE: DNA
    266 <213> ORGANISM: Artificial Sequence
    270 <220> FEATURE:
    272 <223> OTHER INFORMATION: Oligonucleotide
    274 <400> SEQUENCE: 10
    275 catcgatttc ttccctgtga a
                                                                                 21
    278 <210> SEQ ID NO: 11
    280 <211> LENGTH: 24
    282 <212> TYPE: DNA
    284 <213> ORGANISM: Artificial Sequence
    288 <220> FEATURE:
    290 <223> OTHER INFORMATION: Oligonucleotide
    292 <400> SEQUENCE: 11
    293 tcttggagct tattaaaggc attc
                                                                                 24
    296 <210> SEQ ID NO: 12
    298 <211> LENGTH: 24
    300 <212> TYPE: DNA
    302 <213> ORGANISM: Artificial Sequence
    306 <220> FEATURE:
    308 <223> OTHER INFORMATION: Oligonucleotide
    310 <220> FEATURE:
    312 <221> NAME/KEY: misc_feature
    314 <222> LOCATION: (1)..(1)
    316 <223> OTHER INFORMATION ( N \stackrel{\prime}{=} 6Fam
    320 <220> FEATURE:
    322 <221> NAME/KEY: misc_feature
    324 <222> LOCATION: (24)..(24)
    326 <223> OTHER INFORMATION: N = Tamra-p
    330 <400> SEQUENCE: 12
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W--> 331 nacaagagca aggccgtgga gcan

334 <210> SEQ ID NO: 13

24

PATENT APPLICATION: US/10/501,666

DATE: 07/23/2004 TIME: 15:25:18

Input Set : A:\seq\_lst\_US.txt

```
336 <211> LENGTH: 20
              338 <212> TYPE: DNA
              340 <213> ORGANISM: Artificial Sequence
             344 <220> FEATURE:
             346 <223> OTHER INFORMATION: Oligonucleotide
             348 <400> SEQUENCE: 13
             349 tgaggagctg gtcaacatca
                                                                                                                                                                                                  20
             352 <210> SEQ ID NO: 14
             354 <211> LENGTH: 21
             356 <212> TYPE: DNA
             358 <213> ORGANISM: Artificial Sequence
             362 <220> FEATURE:
             364 <223>. OTHER INFORMATION: Oligonucleotide
             366 <400> SEQUENCE: 14
            367 caggttgatg ctccatacca t
                                                                                                                                                                                                  21
             370 <210> SEQ ID NO: 15
             372 <211> LENGTH: 22
            374 <212> TYPE: DNA
            376 <213> ORGANISM: Artificial Sequence
            380 <220> FEATURE:
            382 <223> OTHER INFORMATION: Oligonucleotide
            384 <220> FEATURE:
            386 <221> NAME/KEY: misc_feature
            388 <222> LOCATION: (1)..(1)
            390 <223> OTHER INFORMATION: N = 6FAM
                                                                                                               same
            394 <220> FEATURE:
            396 <221> NAME/KEY: misc_feature
            398 <222> LOCATION: (22)..(22)
            400 <223> OTHER INFORMATION: N = Tamra-p
            404 <400> SEQUENCE: 15
W--> 405 naggeteege tetgeaatgg en
                                                                                                                                                                                                 22
            408 <210> SEQ ID NO: 16
            410 <211> LENGTH: 21
           412 <212> TYPE: DNA
           414 <213> ORGANISM: Artificial Sequence
           418 <220> FEATURE:
           420 <223> OTHER INFORMATION: Oligonucleotide
           422 <400> SEQUENCE: 16
           423 cccagggacc tctctctaat c .
                                                                                                                                                                                                 21
           426 <210> SEQ ID NO: 17
           428 <211> LENGTH: 21
           430 <212> TYPE: DNA
 432 <213> ORGANISM: Artificial Sequence
                                                                                                                                      والمراوية والمرا
           436 <220> FEATURE:
           438 <223> OTHER INFORMATION: Oligonucleotide
           440 <400> SEQUENCE: 17
           441 atgggctaca ggcttgtcac t
                                                                                                                                                                                                21
           444 <210> SEQ ID NO: 18
           446 <211> LENGTH: 24
```

PATENT APPLICATION: US/10/501,666

DATE: 07/23/2004 TIME: 15:25:18

24

24

21

Input Set : A:\seq\_lst\_US.txt

Output Set: N:\CRF4\07232004\J501666.raw

same

same

```
448 <212> TYPE: DNA
```

450 <213> ORGANISM: Artificial Sequence

454 <220> FEATURE:

456 <223> OTHER INFORMATION: Oligonucleotide

458 <220> FEATURE:

460 <221> NAME/KEY: misc\_feature

462 <222> LOCATION: (1)..(1)

464 <223> OTHER INFORMATION: 'N = 6Fam

468 <220> FEATURE:

470 <221> NAME/KEY: misc\_feature

472 <222> LOCATION: (24)..(24)

474 <223 > OTHER INFORMATION: N = Tamra-p

478 <400> SEQUENCE: 18

## W--> 479 ntggcccagg cagtcagatc atcn

482 <210> SEQ ID NO: 19

484 <211> LENGTH: 24

486 <212> TYPE: DNA

488 <213> ORGANISM: Artificial Sequence

492 <220> FEATURE:

494 <223> OTHER INFORMATION: Oligonucleotide

496 <400> SEQUENCE: 19

497 ctaattattc ggtaactgac ttga

500 <210> SEQ ID NO: 20

502 <211> LENGTH: 21

504 <212> TYPE: DNA

506 <213> ORGANISM: Artificial Sequence

510 <220> FEATURE:

512 <223> OTHER INFORMATION: Oligonucleotide

514 <400> SEQUENCE: 20

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518 <210> SEQ ID NO: 21

520 <211> LENGTH: 26

522 <212> TYPE: DNA

524 <213> ORGANISM: Artificial Sequence

528 <220> FEATURE:

530 <223> OTHER INFORMATION: Oligonucleotide

532 <220> FEATURE:

534 <221> NAME/KEY: misc\_feature

536 <222> LOCATION: (1)..(1)

538 <223> OTHER INFORMATION: N = 6Fam

542 <220> FEATURE:

544 <221> NAME/KEY: misc\_feature

546 <222 LOCATION: (26) (26+

548 <223> OTHER INFORMATION: N = Tamra-p

552 <400> SEQUENCE: 21

W--> 553 ntccaacgca aagcaataca tgaacn

556 <210> SEQ ID NO: 22

558 <211> LENGTH: 21

560 <212> TYPE: DNA

Please correct subsequent sequerces 26 containing this error. Sel p. 6 for more error

6

<400> 55 attcagctcg aacactttga a

27 delete at end of file

21

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/501,666

DATE: 07/23/2004 TIME: 15:25:19

FYI

Input Set : A:\seq\_lst US.txt

Output Set: N:\CRF4\07232004\J501666.raw

#### Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:3; N Pos. 1,27
Seq#:6; N Pos. 1,22
Seq#:9; N Pos. 1,29
Seq#:12; N Pos. 1,24
Seq#:15; N Pos. 1,22
Seq#:18; N Pos. 1,24
Seq#:21; N Pos. 1,26
Seq#:24; N Pos. 1,22
Seq#:27; N Pos. 1,31
Seq#:30; N Pos. 22
Seq#:31; N Pos. 1,23
Seq#:48; N Pos. 1,23
Seq#:51; N Pos. 1,23

#### VERIFICATION SUMMARY

j

PATENT APPLICATION: US/10/501,666

DATE: 07/23/2004 TIME: 15:25:19

Input Set : A:\seq lst US.txt

```
L:17 M:270 C: Current Application Number differs, Replaced Application Number.
L:18 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:109 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:3 after pos.:0
L:183 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:6 after pos.:0
L:257 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:9 after pos.:0
L:331 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:12 after pos.:0
L:405 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:15 after pos.:0
L:479 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:18 after pos.:0
L:553 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:21 after pos.:0
L:627 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:24 after pos.:0
L:701 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:27 after pos.:0
L:765 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:30 after pos.:0
L:803 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:31 after pos.:0
L:1129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:48 after pos.:0
L:1203 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:51 after pos.:0
L:1278 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:1280 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1280 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:1282 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1282 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:1284 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
L:1284 M:334 W: (2) Invalid Amino Acid in Coding Region, NUMBER OF INVALID KEYS:1
L:1289 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:55
```

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